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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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       February 13, 2001, 18:08:46; Search time 1088.91 Seconds (without alignments) 12188.502 Million cell updates/sec
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Copyright (c) 1993 - 2000 Compugen Ltd.
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| 190: 191: 192: 193: |
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| gb_gss25:* gb_gss26:* gb_gss27:* gb_gss28:* |

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| | | a | o | | | | | c | C | c | | | | | | c | | C | c | | | | ი | | c | C | | O | c | | n | | c | | a | | | | | | | | | Re |
|---------------|----------------|---------------|--------------|--------------------|------------------|----------|--------------------------|-----------------|-----------|---------------|-------------|--------------|------------|----------|----------|----------|------------------|-----|--------|---------|-------|-----------|----------|------------|----------------|----------|--------|----------|--------|---------------------------------------|----------------|------------|--------------------|-----------------|----------|----------|-----------|--------------|------------------|----------------|------------------|----------------|--------------------|------------------|
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| 277.2 | 279.4 | 279.4 | 280.2 | | \sim | 9 | • | | 312.4 | т- | ٠. | ا ا ا | ٠. | د | 2 | 52. | 357.8 | 59. | 61 | 69. | 69. | 7 | 373 | 76. | • | 8 | 385.8 | 39 | 391.2 | 14 | | 41 | 130 / | 4. | 7 | 488 | 492.8 | ٠ | 518 | 31. | Ģ | 38. | CT. | Score |
| 14.6 | 14.8 | | 14.8 | | 5 | 5 | | 16.2 | 6 | 6 | 6 | ٠,٥ | | 17.6 | • | | 18.9 | 9 | 9. | 9 | 19.5 | 9 | 9 | 9. | .0 | | | | 20.7 | - ! | ٠. | ٥. | ٠. | ٠. | .51 | .51 | | 9 | 7 | 8 | 8 | ٠ | 29.2 | Query Match I |
| 323 | 604 | 285 | 353 | 298 | 318 | 537 | 418 | 364 | 315 | 536 | 600 | 9 9 | 1 (L) | 343 | 369 | 417 | 401 | 563 | 417 | 412 | 496 | 421 | 381 | 518 | 456 | 385 | 471 | 528 | 481 | \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ | 7 = 7 | 7.1 | л 4 л ст 4 | 475 | 602 | 607 | 588 | 681 | 626 | 586 | 682 | 676 | 676 | Length |
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| AW020278 | BF046131 | F03207 | N46450 | 125 | 435 | AI425785 | AW619432 | ω | 396 | T89472 | | AW012792 | | AA307980 | AA308453 | AI493368 | - | | 880 | 2848 | 2729 | W78433 | 88 | BE720244 | 2970 | AA468660 | T74333 | A143352 | N51100 | 22021284 | AA858126 | 012070 | AL924/93 | A1934807 | AA872079 | AV612885 | 78391 | 4 | BF030864 | 9 | 195621 | AI956401 | AI956397 | ID |
| 278 df07d03.y | 46131 BP250004 | 207 HSC1FF112 |) yy76q11.s1 | F12504 HSC3AH031 n | Z43599 HSC1GA021 | 35 me87e | 32 ⁻ 7488 MAR | 9039 yc84f03.s1 | HSC1GA022 | 72 ye04q04.s1 | 364 6012250 | 792 | 64 zr82b09 | EST17882 | EST17927 | 3 tg70g1 | W429512 67771 MA | . ' | E08838 | E284810 | 27291 | 7 zh61d11 | * 104120 | 44 RCO-HTO | 129709 ac35h10 | ne06c01 | T74333 | 72056b11 | | OIGSTO | 5091/9 6010950 | um07e02.y | AI924793 wn23a04.x | 934807 wp89f11. | 9 | ω, 5 | vz75e05.r | 77443 601228 | BF030864 6015576 | 1030 ze46c10 r | 1956216 ul72g10. | 1956401 ul74q1 | AI956397 1174cn6.v | |

ALIGNMENTS

RESULT 1
AI956397
LOCUS
DEFINITION

KCNO1.; mRNA sequence. A1956397 A1956397.1 GI:5749106 EST AI956397 676 bp mRNA EST 20-AUG-1999 ul74g06.yl Sugano mouse kidney mkia Mus musculus cDNA clone IMAGE: 2136346 5', similar to TR:000180 000180 POTASSIUM CHANNEL

ACCESSION VERSION KEYWORDS

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TITLE
JOURNAL
COMMENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     263 GCTGGGCTACTTGCTCTACCTGGTCTTCGGCGCAGTGGTCTTCTCCTCCGTGGAGCTGCC 322
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                                                                                                                                                                                                                                                                                                               GCTCTTCCTTCGCCAGCACCGTGCTCTCCCACCACAGGTTATGGCCACACCGTGCCCTTGTC
                                                                                                                                                                                                                                                                                                                                                                       CTATGAGGACCTGCGCCAGGAGCTGCGCAAGCTGAAGCGACGCTTCTTGGAGGAGCA 382
                                                                                          TTATGGAGTGTCGGTGCTCAGCAACGCCTCGGGAAATTGGAATTGGGACTTCACCTCGGC
                                                                                                                    CTACGGCGTGTCGGTGCTCAGCAACGCCTCGGGCAACTGGA:\CTGGGACTTCACCTCCGC
                                                                                                                                                                                                   CGAGTGCCTGTCGGAGCCGCAGCTGGAGCAGTTCCTGGGCCGCGTGCTGGAGGCCAGCAA
                                                                                                                                                                                                                                                        CGAGTGCCTGTCTGAGCAGCAGCTGGAGCAGTTCCTGGGCCGGGTGCTGGAGGCCAGCAA 442
                                                                                                                                                                                                                                                                                                                                                                                                                              GCTGGGCTACCTTTTCTACCTGGTGTTCGGCGCCGTGGTCT::CTCGTCCGTGGAGCTGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: mouseest@watson.w
This clone is available
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Marra M/WashU-NCI Mouse EST Project 1999 Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Marra,M., Hiller,L., Kucaba,T., Martin,J., Beck,C., Wylle,T., Underwood,K., Steptoe,M., Theising,B., Hallen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Halley,N., Schurk,R., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.
The WashU-NCI Mouse EST Project 1999
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ligated to a DraIII adaptor [IGTTGGCCTACTGG], digested and cloned into distinct DraIII sites of the pME18S-FL3 vector [5, site CACTGTGTG, 3, site CACCATGTG]. MoI should be used to isolate the cDNA insert. Size selection was performed to exclude fragments <1.5kb. Library constructed by Dr. Sumio Suganc (University of Tokyo Institute of Medical Science). Custom primers for sequencing: 5 end primer CTTCTGCTCTAAAAGCTGCG and 3, end primer CGACCTGCAGCTCGAGCACA."

210 c 184 g 173 t 2 others
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/db_xref="taxon:10090"
/clone="IMAGE:2136346"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Organ: kidney; Vector: pME18S-FL3; Site_1: DraIII (CACTGTGTG); Site_2: DraIII (CACTGTGTG); 1st strand cDNA was primed with an oligo(dT) primer with an oligo(dT) primer (ATGTGGCCTTTTTTTTTTTTTTTTTT]; dcuble-stranded cDNA was interested to the strander of the strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Sugano mouse
/sex="female"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 552.4; DB 27; Pred. No. 2.3e-111;
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GAGATTTGTCACCGTTTCCTGCTTCTTCATCCCAGCCGCCGTGTTCTCTGTGCTGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TGGGTTTGTCACTGTGTCCTGCTTCTTCATCCCGGCCGCTGTCTTCTCAGTCCTGGA 802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTTCCTGACGGCTGTGGTCCAGCGCATCACCGTGCACGTCACCCGCAGGCCGGTCCTCTA 682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGATGGAGGTAAGGCCTTCTGCATCTACTCCGTCATTGGCATTCCCTTCACCCTCCT 622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGATGACTGGAACTTCCTCGAGTCTTTTTACTTCTGTTTCATCTCCCTGAGCACCATNCG
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AI956401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AI956401 676 bp mRNA EST 20-AUG-1999 ul74g11.yl Sugano mouse kidney mkia Mus musculus cDNA clone IMAGE:2136356 5' similar to TR:000180 000180 POTASSIUM CHANNEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.
The WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                              Email: mouseest@watson.wustl.edu
This clone is available royalty-free
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI956401.1 GI:5749110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Marra M/WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                         MGI:100103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                            Possible reversed clone: similarity
                                                                                                                                                                                                                                                                                                                                                                    IMAGE Consortium (info@image.llnl.gov)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mmalia; Eutheria; (bases 1 to 676)
                                                                                                                                                                                                                                                                                  primer: custom primer used
                                                                                                                                                                                                                             quality sequence stop:
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;, mRNA sequence.
  /sex="female"
/dev_stage="adult"
                                                       /clone="IMAGE:2136356"
/clone_lib="Sugano mouse kidney mkia"
                                                                                                                 /strain="C57BL"
/db_xref="taxon:10090"
                                                                                                                                                                      /organism="Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     676
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                                                                                                                                                                                                                                                                                                                                                                    through LLNL;
v) for further
                                                                                                                                                                                                                                                                                                                                                                          further information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             63108, USA
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660 922 600 862 540 480

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Query Match
Best Local Similarity
Matches 578; Conserv
                                                                                                                                                                                                                                                                                                                                                                                     GTCCTGCTTCTTCATCCCGGCCGCTGTCTTCTCAGTCCTGGAG 803
                                                                                                                                                                                                                                                                                                                                     GGTCCAGCGCATCACCGTGCACGTCACCCGCAGGCCGGTCCTCTACTTCCACATCCGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCGCCAGGAGCTGCGCAAGCTGAAGCGACGCTTCTTGGAGGAGCACGAGTGCCTGTCTGA 397
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                                                                                                                                                                                                                                     NGGCTTCTCCAAGCAGGTGGTGGCCATTGTCCATGCCGTTCTGNCTGGATTTGTCACCGT
                                                                                                                                                                                                                                                                     GGGCTTCTCCAAGCAGGTGGTGGCCATCGTCCATGCCGTGCTCCTTGGGTTTGTCACTGT 757
                                                                                                                                                                                                                                                                                                               GGTCCAGCGTGTCACCGTGCATGTCACCCGCAGACCAGTCCTCTACTTCCACATACGCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTCCTGCTTCTTCATCCCCAGCCGCGTGTTCTCTGTGCTGGAG
                A1956216 682 bp mRNA EST u172g10.yl Sugano mouse kidney mkia Mus muscu IMAGE:2136162 5' similar to TR:000180 000180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="DH108"
//note="Organ: kidney; Vector: pME18S-FL3; Site_1: DraIII
//note="Organ: kidney: DraIII (CACCATGTG); 1st strand cDNA
was primed with an oligo(dT) primer
[ATGTGCCTTTTTTTTTTTTTTTTT]; double-stranded cDNA was
ligated to a DraIII adaptor [TGTTGCCTACTGG] cDNA was
ligated to a DraIII adaptor [TGTTGCCTACTGG]. XhoI should
and cloned into distinct DraIII sites of the pME18S-FL3
vector (5' site CACTGTGTG, 3' site CACCATGTG). XhoI should
be used to isolate the cDNA insert. Size selection was
performed to exclude fragments <1.5kb. Library
constructed by Dr. Sumio Sugano (University of Tokyo
Institute of Medical Science). Custom primers for
sequencing: 5' end primer CTGTCTGTCTTAAAAGCTGCG and 3' end
primer CGACCTGCAGCACACA."

94 a 216 c 211 g 153 t 2 others
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    ;, mRNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 538.4; DB 27; Pred. No. 2.8e-108;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                    musculus cDNA clone
00180 POTASSIUM CHANNEI
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                                                            20-AUG-1999
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ORGANISM
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217
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                                                                                                                                                                                            218
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                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                              GCGCCAGGAGCTGCCGCAAGCTGAAGCGACGCTTCTTGGAGGAGCACGAGTGCCTGTCTGA
                                                                                                                                                                          CCTGGTGGAGCGGCACCGCTCGGCCTGGTGCTTCGGCTTCCTGGTGCTGGTGCTTGGT 277
                                                                                              CTACCTGGTCTTCGGCGCAGTGGTCTTCTCCTCGGTGGAGCTGCCCTATGAGGACCTGCT
                                                                            CTACCTGGTGTTCGGCCGTGGTCTTCTCGTCCGTGGAGCTGCCTTATCAGGACCTGCT
                                                                                                                                                      CCTGGTGGAGCGGCACCGTTCGGCCTGGTGCTTCCGGCTTCCTGGTGCTGGCCTACCTGCT
GCGCCAGGAGCTGCGCAAGCTGAAGCGGCGCTTCCTGGAGGAGCACGAGTGCCTGTCGGA
                                                                                                                                                                                                                                                                                                             576;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: mouseestéwatson.wustl.edu
Email: mouseestéwatson.wustl.edu
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.
The WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Marra M/WashU-NCI Mouse EST Project 1999 Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Possible reversed clone: similarity Seq primer: custom primer used High quality sequence stop: 503.
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AI956216.1
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Mammalia; Eutheria;
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314 286 1810
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216 c 215 g 154 t 2 others
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/clone="IMAGE:2136162"
/clone_1lb="Sugano mouse kidney mkia"
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4444 Forest Park Parkway,
Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 586)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lenncn,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.

The WashU-Merck EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA001030 586 bp mRNA EST 29-NO
ze46c10.rl Soares retina N2b4HR Homo sapiens cDNA clone
IMAGE:362034 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1995)
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                                                                                                                                                                                                                                                                                                                                                                  Contact: Wilson RK
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                                                                                                                                                                                                        quality sequence stop: 445.
/tissue_type="retina"
/dev_stage="55 year old"
/lab_host="0H10B (ampicillin resistant)"
/note="Organ: eye; Vector: pT7F3D (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; :
                                                                                           /organism="Homo sapiens"
/db_xref="GDB:1278737"
/db_xref="taxon:9606"
/clone="IMAGE:362034"
/clone=lib="Soares retina N2b4HR"
                                                                                /sex="male"
                                                                                                                                                                                          ocation/Qualifiers
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Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 626)
                                                                                                                                            mRNA sequence.
                                                                                                                                                               BF030864 626 bp mRNA EST 10-OCT-2000 601557647F1 NIH_MGC_58 Homo sapiens cDNA clone IMAGE:3827531
                                                    Homo sapiens
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98.3%;
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                                                                                                                                                                                                                                    1141 CTTTTGTGGCCACCCAGTCATCTGCCTGCGTGGATGGCCCTGCAAACCATTGAGCGTAGG 1200
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                    ATCTCGACCTTACATAGGAGGAGAATACTTGAAGCAGTATGCTGCTGTGGTTAGAAGCAG 1499
                                                                                                                                                                                AATGTCTTATTAAAAAACAAC-AAAAAAAGGACACATGGAACAAGGAAGGTGTGACCCCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CCTTCTCCTCGATCACAGACCAGGCAGCTGGCATGAAAGGAGGACCAGAAGCAAAATGAGC 120
ATCTCGACCTTACATAGGAGGAGAATACTTGAAGCAGTATGCTGCTGTGGTTAGAAGCAG
                                                                            CAGGATGTCTAATATGTGAGGAAATGAGATGTCCACCTAAAATTCATATGTGACAAAATT
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                                                                                                                                                            AATGTCTTATTAAAAAACAACAAAAAAAAAAGACAAATGGAACAAAGAAGCTGTGACCCCAG
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Email: Robert_Strausberg@nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     http://image.llnl.gov
Plate: LLCM496 row: n column: 12
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National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              215 a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="hypernephroma"
/tlssue_type="hypernephroma"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: kidney; Vector: pDNR-LIB (Clontech); Site_1:
Sii (ggccgctctggcc); Site_2: Sfii (ggccattatggcc);
Double-stranded cDNA was prepared from cell line RNA. 5'
and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAGAGGCCGAGAGG-GT(30)BN-3'
(where B = A, C, or G and N = A, C, G, or T). Average
insert size 1.35 kb (range 0.9-4.0 kb). 15/15 colonies
contained inserts by PCR. This library was enriched for
full-length clones and was constructed by Clontech
Laboratories (Palo Alto, CA)."
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/clone_lib="NIH_MGC_58"
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/db_xref="taxon:9606"
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                                                 CCTGGTGGAGCGGCACCGCTCGGCCTGGTGCTTCGGCCTTCCTGGTGCTGGGCTACTTGCT 277
CTACCTGGTCTTCGGCGCAGTGGTCTTCTCCTCGGTGGAGCTGCCCTATGAGGACCTGCT 337
                             CCTGGTGGAGCGGCACCGTTCGGCCTGGTGCTTCGGCTTCCTGGTGCTGCGCTACCTGCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM8764 row: h column: 05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: Robert_Strausberg@nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortiu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 599.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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                                                                                                                                                                                                                                                                     /note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: Site_2: NotI; Cloned unidirectionally. Primer: Oilg Library constructed by Life Technologies. Investigat providing samples: Gilbert Smith, NIH"

216 c 213 g 158 t
                                                                                                                                                                                                                                                                                                                                                     /clone="IMAGE:3592756"
/clone_lib="WCI_CGAP_Mam1"
/tissue_type="tumor, biopsy sample"
/dev_stage="3 months, virgin"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /strain="FVB/N"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Mus musculus"
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88.7%;
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Pred. No. 2.7
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IMAGE:3592756 5',
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                                                                                                                                                                                              Unpublished (195)
Unpublished (195)
Contact: Wilson RK
Washington University School of Medicine
Washington University School of Medicine
                                                                                    Seq primer: reverse in 394 High quality sequence stop: 394 Location/Qualifiers
                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 588)

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennen, G., Marra, M., Parsons, J., M., Hultman, M., Kucaba, T., Le, M., Lennen, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.

The WashU-Merck EST Project
                                                                                                                                                                         4444 Forest Park Parkway,
Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                              N78391 588 bp mRNA EST yz75e05.rl Soares_multiple_sclerosis_2NtHMSP clone IMAGE:288896 5', mRNA sequence.
                                                                                                                                               Email: est@watson.wustl.edu
This clone is available roy
                                                                                                                                                                                                                                                                                                                                                                                                        N78391.1
                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                    IMAGE Consortium (info@image.llnl
/organism="Homo sapiens"
/db_xref="GDB:3904736"
/db_xref="taxon:9606"
/clone="IMAGE:288896"
/clone_lib="Soares_multiple_sclerosis_2NbHMSP"
/sex="male"
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GACGAGGATCAGGTGCACATCATAGAGCATGACCAACTGTCCTTCTCCTT-CGATCACAGA 1099
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                                                                                                                                                                                                                                                     GAGCACCAGGGTCAGG 1234
                                                                                                                                                                                                                                                                                               CATCTGCCTGCGTGGATGGGCCTGCAAACCATTGAGCGTAGGANTTGGTGCATTATNCTA 542
                                                                                                                                                                                                                                                                                                                                                                             CCAGGCAGCTGGGCATGAAAGAGGACCAGAAGCAAATGAGCCTTTTGGTGGCCACCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                      CCAGGCAGCT-GGCATGAAAGAGGACCAGAAGCAAAATGAGCCTTTTGTGGCCACCCAGT 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                GACGAGGATCAGGTGCACATCATAGAGCATGACCAACTGTCCTTCTCCTCCGATCACAGA
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AV612885 BOS ta

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                                                                                  Bos taurus lung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="multiple sclerosis lesions"
/dev_stage="Age 46"
/lab_host="NH10B (ampicillin resistant)"
/note="Vector: pT7T3D (Pharmacia) with a modified polylinker V_TYPE: phagemid; Site_1: Not I; Site_2: E; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from 4 multiple sclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lesions from one patient was kindly provided by Dr. Kevin
G. Becker (NINDS/NIH). "
134 c 154 g 160 t 4 others
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492 GGTCTCATTGCAATGTTGGTGGTTCTTGGAAACCTTCTGTGAGCTCCACGAACTGAAGAAG
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                                                                                                                                                                                                                                                                                                                                                   192 GTTACCATCCATGTCACCCGCAGGCCAGTCCTCTACTTCCACGTCCGCTGGGGTTTCTCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72 TCCACCACAGGTTACGGCCACACCGTGCCCCTGTCCGACGGGGGCAAGGCCTTCTGCATC
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                                                                                                                                                                                                                                                                                                                                                                         ATCACCGTGCACGTCACCCGCAGGCCGGTCCTCTACTTCCACATCCGCTGGGGCTTCTCCC
                                GGCCTTATTGCCATGTTGGTAGTTCTGGAAACCTTCTGTGAACTCCATGAGCTGAAAAAA
                                                                                     GGCTACAATCAAAATTCAGAGAGCTCTATAAGATTGGGATCACGTGTTACCTGCTACTT
                                                                                                                                                          TTTTATTTTTGTTTTATTTCCCTGAGCACCATTGGCCTGGGGGATTATGTGCCTGGGGAA
                                                                                                                                                                                                                                                                                AAGCAGGCAGTGGCCATCGTCCACGCGGTCCTCCTCGGTGTGGTCACCGTGTCCTTC
                                                                    GGTTACAATCAAAAATTCAGAGAGCTCTACAAGATCGGGATCACATGTTACCTGCTCCTT
                                                                                                                                        TTTTACTTTTGTTTTATCTCCCTGAGCACCATTGGCCTCGGAGATTATGTTCCCGGGGAG
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Shirakawa Institute of Animal Genetics
Odakura, Nishigo, Nishi-shirakawa, Fuku
Tel: 81-248-25-5641
Fax: 81-248-25-5725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: kazusugi@cocoa.ocn.ne.jp
Single pass sequencing.
This clone was obtained from a polyA-deleted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cow.
Bos taurus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Yoshikazu Sugimoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bovine cDNA sequencing Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sugimoto,Y., Hirotsune,S., Suzuki,H.
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was deleted from a Not1 site"
184 c 148 g 157 t
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/dev_stage="fetus"
/lab_host="DH10B"
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/db_xref="taxon:9913"
/clone="E1LU048D11"
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TITLE
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Best Local Similarity
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                     1038 AAGGACGAGGATCACGTGCACATCATAGAGCATGACCAACTGTCCTTCTCCTCCGATCACA 1097
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GACGGCGATGATCAGGTGCACATCATAGAGCATGACCAACTGTCCTCTCCCTCGATAACA
                                                                                                ACCTTCTGTGAACTCCATGAGCTGAAAAAATTCAGAAAAATGTTCTATGTGAAGAAGACAC 1037
                                                                             ACCCTCTGTCGACTCCATGAGCT--AAAAAATCAGAATAATG-TCTATGTGAAG-ACGAC 547
                                                                                                                                                                 552;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortlum/LLNL at: www-bio.llnl.gov/bbrp/image/image.html Seg primer: -40ml3 fwd. ET from Amersham
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oil2a07.sl
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Tissue Procurement: Christopher A. Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 004)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:1476276"
/clone_lib="NCI_CGAP_GC4"
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1098 GACCAGGCAGGCATGAAAGAGGACCAGAAGCAAAATGAGCCTTTTGTGGCCACCCAG 1157

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1 (bases 1 to 475)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gcv/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project
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wp89f11.x1
                                                                                                                                    DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
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                                                                                                      quality sequence stop: 444.
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                                                                                                                                                                                                                                                                                   Robert_Strausberg@nih.gov
Procurement: David N. Louis, M.D., Myrna R.
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2468973"
/clone_lib="NCI_CGAP_Brn25"
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                                                                                                                                                                                                             mRNA sequence.
AI924793
AI924793.1 GI:5660757
                                             Unpublished (1997)
Contact: Robert Strausberg,
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wn23a04.x1 NCI_CGAP_Gas4
                               Tel: (301) 496-1550
                                                                            Tumor Gene Index
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1 (bases 1 to 484)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Ana
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Robert_Strausberg@nih.gov
Procurement: Christopher
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/lab_host="DH10B"
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Moskaluk,

M.D.,

Ph.D.,

Michael

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ACCESSION
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                                                                                                                                  RESULT 1
AW012076
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                                                                          AW012076 551 bp mRNA EST 10-SEP-19-
um07e02.y1 Sugano mouse kidney mkia Mus musculus cDNA clone
IMAGE:2182394 5' similar to TR:008581 008581 POTASSIUM
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High quality sequence stop: 423.
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Insert Length: 1419 Std Error: 0.00
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AW012076
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                                                           INWARDLY-RECTIFYING
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Site_2: NotI; Cloned unidirectionally. Primer:
Average insert size 1.69 kb. Life Technologies
11549-011"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              signet ring
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/clone_lib="NCI_CGAP_Gas4"
/tissue_type="poorly differentiated adenocarcinoma
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/lab_host="DH10B"
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                                                                                                                                                                                                                                                                  CTACCTGGTCTTCGGCGCAGTGGTCTTCTCCTCGGTGGAGCTGCCCTATGAGGACCTGCT 337
                              GCGCCAGGAGCTGCGCAAGCTGAAGCGGCGCTTCCTGGAGGAGCACGAGTGCCTGTCGGA
                                                                                                                                                                 GCGCCAGGAGCTGCGCAAGCTGAAGCGACGCTTCTTGGAGGAGCACGAGTGCCTGTCTGA 397
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Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.

The WashU-NCI Mouse EST Project 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800
Fax: 314 286 1810
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4444 Forest Park Parkway, Box 8501, St. Louis,
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Contact: Marra M/WashU-NCI Mouse EST Project 1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     constructed by Dr. Sumio Sugano (University of Tokyo Institute of Medical Science). Custom primers for sequencing: 5' end primer CTTCTGCTCTAAAAGCTGCG and 3' primer CGACCTGCAGCTCGAGCACA."
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//lab_host="DH10B"
//note="Organ: kidney; Vector: pME18S-FL3; Site_1: DraIII
//note="Organ: kidney; Vector: pME18S-FL3; Site_1: DraIII
//note="Organ: kidney; Vector: pME18S-FL3; Site_1: DraIII
//note="Organ: kidney; Vector: primer
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/strain="C57BL"
/db_xref="taxon:10090"
/clone="IMAGE:2182394"
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                                                                                                                                                                                                                                                                                                                          DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov h column: 11
Plate: LLAM8530 row: h column: 11
High quality sequence stop: 541.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                   Email: Robert_Strausberg@nih.gov
Tissue Procurement: Lothar Hennighausen Ph.D., Robin
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.3. Consortium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
Contact: Robert Strausberg, Ph.D. Tel: (301) 496-1550
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NIH-MGC http://www.ncbi.nlm.nih.gov/MGC/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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601095044F1 NCI_CGAP_Mam5 Mu:
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                                                                                                                                                      /note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sall Site_2: NotI; Cloned unidirectionally. Primer: Oligo dy Library constructed by Life Technologies. Investigators providing samples: Lothar Hennighausen/Robin Humphreys,
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/dev_stage="7 months"
/lab_host="DH10B"
                                                                                                                                                                                                                                                             /strain="C57/B6"
/db_xref="taxon:10090"
/clone="IMAGE:3489466"
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                                                                                                        CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html Insert Length: 1249 Std Error: 0.00
                                                                                                                                                                                                                                                                         Email: Robert_Strausberg@nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D Tel: (301) 496-1550
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             Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry, Meg, J., Trevaskis, E., Schellenberg, K., Soares, M.B., Tan, F., Thierry, Meg, J., Trevaskis, E.,
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ze68g04.rl Soares retina N2b4HR Homo sapiens cDNA clone
IMAGE:364182 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
/clone="IMAGE:1435261"
/clone_lib="NCI_CGAP_Co8"
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/lab_host="DH10B"
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Wohldmann, P.,
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Pred. No. 1.3e-81;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 12;
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GCTTCTTCATCCCGGCCGCTGTCTTCTCAGTCCTGGAGGATGACTGGAACTTCCTGG
                                                                                   GGGAAGGCTACAATCAAAAATTCAGAGAGCTCTATAAGATTGGGATCACGTGTTACCTGC 942
                                                                                                               AATCCTTTTATTTTTGTTTTATTTCCCTGAGCACCATTGGCCTGGGGGGATTATGTGCCTG
                                                                                                                              AATCCTTTTATTTTGTTTTATTTCCCTGAGCACCATTGGCCTGGGGGGATTATGTGCCTG
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                                                                                                                                                                                                                                             TCTCCAAGCAGGTGGTGGCCATCGTCCATGCCGTGCTCCTTGGGTTTTGTCACTGTGTCCT 762
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                                                        GGGAAGGCTACAATCAAAAATTCAGAGAGCTCTATAAGATTGGGATCACGTGTTACCTGC
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97044478
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Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4444 Forest Park Parkway, Box 8501, St. Tel: 314 286 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73
                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Soares and M.Fatima Bonaldo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="retina"
/dev_stage="55 year old"
/lab_host="DH10B (ampicillin_resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="GDB:1280885"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Soares retina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:364182"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                  21.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of Toronto, Library constructed by Bento M Fatima Ronaldo. "
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Pred. No. 5.2e-81;
0; Mismatches 6
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